

**AMENDMENTS TO THE SPECIFICATION:**

Please replace the paragraph at 18, line 18 to page 19, line 7 with the following revised paragraph:

--A density related to the presence of the peptide could be located after several rounds of refinement in a "simulated annealing composite omit map" (Brunger *et al.*, 1998). The seven C-terminal residues of the P16 peptide, R<sub>10</sub>Q<sub>11</sub>L<sub>12</sub>V<sub>13</sub>L<sub>14</sub>G<sub>15</sub>L<sub>16</sub> (SEQ ID NO: 6) encompassing the  $\beta$  binding sequence were built into the density map (**Figure 2**). This map extended slightly toward the N-terminus of the peptide but rapidly faded, so that the Q<sub>11</sub> residue was still easily seen while the R<sub>10</sub> was built in a poor density region. The rest of the peptide, probably disordered, was not visible. The seven C-terminal amino acids of the P16 peptide bind onto the  $\beta$  surface within two distinct but adjacent domains: one deep crevice, located between sub-domains 2 and 3 (area 1), and a second area which extends over the third  $\beta$  subdomain, close to the C-terminal extremity of the  $\beta$  chain (area 2) (**Figure 2**). --

**In the Abstract:**

Please cancel the originally-filed Abstract of the Disclosure, and add the accompanying new Abstract of the Disclosure which appears on a separate sheet in the Appendix.